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		032301SigM.seq.ST25.txt													
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Ala Phe Ser Thr Ile Ile His Arg His Glu Arg His Met Met Gln Ala 50 60

Ala Arg Lys Tyr Gly Arg Lys Pro Glu Asp Ala Gln Asp Ile Leu Gln 65 70 75 80

Glu Ala Leu Phe Arg Ala Ser Arg Asn Met His Leu Tyr Arg Ala Glu 85 90 95

Ala Ala Leu Gly Thr Trp Leu His Lys Leu Val Leu Asn Ser Gly Phe 100 105 110

Asp Trp Ala Thr His Arg Ser Gln Val Glu Phe Pro Ile Leu Asn Glu 115 120 125

Pro Thr Ile Asp Leu Glu Lys Asp Pro Arg Leu Ala Thr Asp Pro Leu 130 135 140

Gly Tyr Leu Asp Val Ala Met Thr Ile Arg Ser Ala Ile Asp Gln Leu 145 150 155 160

His Pro Asp Gln Arg Ile Ala Leu Ile Leu Val Asp Leu Gly Gly Tyr \$165\$ \$170\$ \$175\$

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<213> Corynebacterium glutamicum

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